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STAZEWSKI, LENA  
XU, HONG  
EHEVERRI, FERNANDO

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Val	Arg	Tyr	Gln	Gly	Pro	Tyr	Val	Ser	Met	Ala	Phe	Ile	Thr	Val	Leu	675	680	685
Lys	Met	Val	Ile	Val	Val	Ile	Gly	Met	Leu	Ala	Thr	Gly	Leu	Ser	Pro	690	695	700
Thr	Thr	Arg	Thr	Asp	Pro	Asp	Asp	Pro	Lys	Ile	Thr	Ile	Val	Ser	Cys	705	710	715
Asn	Pro	Asn	Tyr	Arg	Asn	Ser	Leu	Leu	Phe	Asn	Thr	Ser	Leu	Asp	Leu	725	730	735
Leu	Leu	Ser	Val	Val	Gly	Phe	Ser	Phe	Ala	Tyr	Met	Gly	Lys	Glu	Leu	740	745	750
Pro	Thr	Asn	Tyr	Asn	Glu	Ala	Lys	Phe	Ile	Thr	Leu	Ser	Met	Thr	Phe	755	760	765
Tyr	Phe	Thr	Ser	Ser	Val	Ser	Leu	Cys	Thr	Phe	Met	Ser	Ala	Tyr	Ser	770	775	780
Gly	Val	Leu	Val	Thr	Ile	Val	Asp	Leu	Leu	Val	Thr	Val	Leu	Asn	Leu	785	790	795
Leu	Ala	Ile	Ser	Leu	Gly	Tyr	Phe	Gly	Pro	Lys	Cys	Tyr	Met	Ile	Leu	805	810	815
Phe	Tyr	Pro	Glu	Arg	Asn	Thr	Pro	Ala	Tyr	Phe	Asn	Ser	Met	Ile	Gln	820	825	830
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<211> 2559

<212> DNA

<213> Homo sapiens



<400> 5

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<211> 852

<212> PRT

<213> Homo sapiens

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Ser Gly Cys Leu Arg Gly Pro Trp Ala Trp Leu Val Val Leu Leu Ala	675	680	685			
Met Leu Val Glu Val Ala Leu Cys Thr Trp Tyr Leu Val Ala Phe Pro	690	695	700			
Pro Glu Val Val Thr Asp Trp His Met Leu Pro Thr Glu Ala Leu Val	705	710	715			720
His Cys Arg Thr Arg Ser Trp Val Ser Phe Gly Leu Ala His Ala Thr	725	730	735			
Asn Ala Thr Leu Ala Phe Leu Cys Phe Leu Gly Thr Phe Leu Val Arg	740	745	750			
Ser Gln Pro Gly Cys Tyr Asn Arg Ala Arg Gly Leu Thr Phe Ala Met	755	760	765			
Leu Ala Tyr Phe Ile Thr Trp Val Ser Phe Val Pro Leu Leu Ala Asn	770	775	780			
Val Gln Val Val Leu Arg Pro Ala Val Gln Met Gly Ala Leu Leu Leu	785	790	795			800
Cys Val Leu Gly Ile Leu Ala Ala Phe His Leu Pro Arg Cys Tyr Leu	805	810	815			
Leu Met Arg Gln Pro Gly Leu Asn Thr Pro Glu Phe Phe Leu Gly Gly	820	825	830			
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 <212> DNA  
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<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Consensus sequence

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<223> Thr or Arg

<220>

<221> MOD\_RES

<222> (3)

<223> Phe or Leu



<220>  
<221> MOD\_RES  
<222> (4)  
<223> Arg, Gln or Pro

<220>  
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<222> (6)  
<223> Arg or Thr

<220>  
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<222> (7)  
<223> Ser, Pro or Val

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<213> Artificial Sequence

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<223> Description of Artificial Sequence: Consensus  
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<220>  
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<222> (3)  
<223> Glu, Gly or Thr

<220>  
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<220>  
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<222> (7)  
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<223> Cys, Gly or Phe

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<220>  
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<223> Description of Artificial Sequence: Synthetic  
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